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January 16, 2003, 16:55:57; Search time 440.214 Seconds (Without alignments) 330.553 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Re.R. and Cook.J.
Inhibition of cellular proliferation in vitro by oligonucleotide binding to a chicknesomal binding site for p53 protein Patent: US 6303328-A 9 16-0CF-2001;
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Mulshine,J.L. and Tockman,M.S.
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Mulshine, J.L. and Tockman, M.S.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primales, Catarrhini, Hominidae, Homo.
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obesity associated biallelic marker maps
Patent: Wo 0206525-A 106 24-JAN-2002;
GENSET (FR)
                                                                  Essential genes of yeast as targets for antifungal agents, herbicides, insecticides and anti-proliferative drugs Fatent: US 6200803.A 47 13.MAR-2001,
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                                                                                                                                       /organism-"unknown"
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AX378317.1 GI:19574167
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Roberts, C.J.
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S44200 60 bp mRNA linear PLN 08-MAY-1993 class VI zygote specific cell wall hydroxyproline-rich glycoprotein [Chlamydomonas reinhardtii, mRNA Partial, 60 nt].
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Chlamydomonas reinhardiii.
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Chlamydomonas reinhardiii.
Chlamydomonadaccae. Chlamydomonas.
Chlamydomonadaccae. Chlamydomonas.
I to 60)
Woessner.i.p. and Goodconough, U.W.
Moleviiir characterization of a zygote wall protein: an extensin-like molecule in chlamydomonas reinhardiii
Piant cell 1 (9), 901-911 (1989)
                                                                                                              Method for determining ancestral habiotypes using haplospecific geometric elements within the major histocompatibility complex
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Mismatches:
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Dawkins, R. Letts. and Abraham, L. Joseph.
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Location/Qualifiers
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/db_xref="G1:255404"
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AF062766
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Rosidae; curosids 1; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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Direct Submission
Submitted (04 MAY 1998) Division of Relany and Sectory, Australian
                                                                                                                                                                                                                                                                                                                                Bukanyota: Viridiplantae: Streptophyta: Embryophyta: Tracheophyta;
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Peakall.R., Gilmore,S., Keys,W., Morgante,M. and Rafalski,A.
Cross-species amplification of soybean (Glycine max) simple
sequence repeats (SSRs) within the genus and other legume genera:
implications for the transferability of SSRs in plants
Mol. Biol. Evol. 15 (10), 1275-1287 (1998)
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STS 04-AUG-1994
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Cetartiodactyla, Ruminantia, Pecora, Cervoidea,
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STS, PCR primer, STS sequence, microsatellite DNA; microsatellite
arker, repeat polymorphism; sequence tagged site,
Homo sapiens DNA
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Submitted by: Utah Center for Human Genome Research University of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                       [ (bases 1 to 62)
Fichel,J. and Reinsch,A.
Microsatellite sequences as genetic markers in European Roe deer
                                                                                                                                                                                                                                                             Submitted (02-30L-1999) Evolutionary Genetics, Institute for Zoo Biology and Wildlife Research, Alfred Kewalke-Str. 17, Herlin D-10315, Germany
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Gerken, S.C., Matsunami, N., Plaetke, R., Albertsen, H., Ballard, L.,
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Human SIS UI272, 5' primer bind, sequence tagged sile.
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/db_xret="taxon:9858"
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2160 Eccles Institute of Human Genetics
Salt Lake City, UT 84112
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                                                      Cervidae; Odocoileinae; Capreolus.
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/rpt_type-tandem
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/rpt_unit-qt
31...38
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Fickel, J. and Reinsch, A.
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Saccharomycetales, mitospoile Saccharomycetales; Candida.
                                                                          Cycles Denaturation Annealing Extension 35 C 10 sec. 65 C 10 sec 72 C 20 sec. Mg··. 2.09 mM Gel: Acrylamide 78, Formamide 32%, Urea 34%
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DEFINITION M.musculus microsatellite DNA DONGS26.
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/db_xret-"taxon:5476"
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/db_xrof-"ravon.9κθέ"
/map="17"
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                       TGCAGATACTGGGGGGCCACAGA
                                                               Initial Denaturation: 94C 360sec
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Edwarptu, Mctaroa, Chordata, Cramiata, Vortebrata, Euterioostomi; Mammadia, Eutheria, Rodentia; Sciurognathi; Muridae; Murinae; Mus. (bases 1 to 67)
Cornall, E.J., Attman, T.J., Hearne, C.M. and Todd, J.A. The generation of a library of Por-analyzed microsatellite variants for quentic mapping of the mouse genome
                                                                                                                                                                                                                                                             2 (bases 1 to 67)
Cornell, R.C., Altman, T.J., Hearne, C.M. and Todd, J.A.
The Generation of a Library of PCR-Analysed Microsatellite Variants
for Genetic Mapping of the Mouse Geneme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Inokuchi,Y., Hirashima,A. and Watanabe,I.
Comparison of the nucleotide sequences at the 3' terminal region of SNAs from RNA coliphages
J. Moi. Biol. 158 (4), 711-730 (1982)
7120417
                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (22-oct-1940) C.M. Hearne, UNIVERSITY OF OXFORD, NUFFIELD DEPT. SURGERY, LEVEL 6, JOHN RADCLIFFE HOSP., HEADINGTON OXFORD, UK
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Allokevivirus; Allokevivirus subgroup IV.
1 (bases 1 to 78)
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Mismatches:
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/db_xref="taxon:10090"
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/organism-"Mus musculus"
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linear PAT 25-SEP-2001
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synthetic construct
artificial sequences.
1 (bases 1 to 83)
Roemer,T., Jianq,B., Boone,C. and Bussey,H.
Gene disruption methodologies for drug target discovery
Patent: Wo 0160975-A 218 23-AuG-2001;
Elitra Pharmacenticals, Inc. (US)
         78
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